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Sequence Listing was accepted.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)

217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: [year=2008; month=8; day=6; hr=15; min=6; sec=18; ms=991; ]

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## Validated By CRFValidator v 1.0.3

Application No: 10565595 Version No: 1.0

Input Set:

Output Set:

**Started:** 2008-06-30 13:47:05.063

**Finished:** 2008-06-30 13:47:06.376

Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 313 ms

Total Warnings: 9

Total Errors: 5

No. of SeqIDs Defined: 9

Actual SeqID Count: 9

Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (1)
W 213	Artificial or Unknown found in <213> in SEQ ID (2)
W 213	Artificial or Unknown found in <213> in SEQ ID (3)
W 213	Artificial or Unknown found in <213> in SEQ ID (4)
W 213	Artificial or Unknown found in <213> in SEQ ID (5)
E 257	Invalid sequence data feature in <221> in SEQ ID (5)
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W 213	Artificial or Unknown found in <213> in SEQ ID (6)
W 213	Artificial or Unknown found in <213> in SEQ ID (7)
W 213	Artificial or Unknown found in <213> in SEQ ID (8)
W 213	Artificial or Unknown found in <213> in SEQ ID (9)

## SEQUENCE LISTING

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<110> ARAKAWA, TAKESHI
     KIKUKAWA, MASANAO
      SHIMABUKURO, ISAO
      TADANO, MASAYUKI
      MATSUMOTO, YASUNOBU
      TSUJI, NAOTOSHI
      SATO, YOSHIYA
<120> HETERO TYPE PENTAMER RECOMBINANT VACCINE
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<140> 10565595
<141> 2008-06-30
<150> PCT/JP04/10459
<151> 2004-07-23
<150> JP 2003-279156
<151> 2003-07-24
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Tyr His Asn Thr Gln Ile His Thr Leu Asn Asp Lys Ile Phe Ser Tyr
        35
                            40
                                                 45
Thr Glu Ser Leu Ala Gly Lys Arg Glu Met Ala Ile Ile Thr Phe Lys
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                         55
Asn Gly Ala Thr Phe Gln Val Glu Val Pro Gly Ser Gln His Ile Asp
                     70
                                         75
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Ser Gln Lys Lys Ala Ile Glu Arg Met Lys Asp Thr Leu Arg Ile Ala

85 90 95

Tyr Leu Thr Glu Ala Lys Val Glu Lys Leu Cys Val Trp Asn Asn Lys
100 105 110

Thr Pro His Ala Ile Ala Ala Ile Ser Met Ala Asn Gly Pro Gly Pro 115 120 125

Glu Phe Thr Tyr Gly Met Cys Thr Glu Lys Phe Ser Phe Ala Lys Asn 130 135 140

Pro Ala Asp Thr Gly His Gly Thr Val Val Ile Glu Leu Ser Tyr Ser 145 150 155 160

Gly Ser Asp Gly Pro Cys Lys Ile Pro Ile Val Ser Val Ala Ser Leu 165 170 175

Asn Asp Met Thr Pro Val Gly Arg Leu Val Thr Val Asn Pro Phe Val
180 185 190

Ala Thr Ser Ser Ala Asn Ser Lys Val Leu Val Glu Met Glu Pro Pro 195 200 205

Phe Gly Asp Ser Tyr Ile Val Val Gly Arg Gly Asp Lys Gln Ile Asn 210 215 220

His His Trp His Lys Ala Gly Ser Thr Leu Gly Lys Ala Phe Ser Thr 225 230 235 240

Thr Leu Lys Gly Ala Gln Arg Leu Ala Ala Leu Gly Asp Thr Ala Trp
245 250 255

Asp Phe Gly Ser Ile Gly Gly Val Phe Asn Ser Ile Gly Lys Ala Val 260 265 270

His Gln Val Phe Gly Gly Ala Phe Arg Thr Leu Phe Gly Gly Met Ser 275 280 285

Trp Ile Thr Gln Gly 290

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